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sequence, subsequence display, backbone, repeat region

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### WD Repeat example

Each repeat **sequence** is broken into columns that describe the structure of the ... The conserved portions of the WD **repeat region** itself were shadowed with X's ... The BLASTP search results are as follows: \* The amino **subsequence** is too .... The model coordinates are for the **backbone** atoms and beta carbons only of ...

bmrcr-www.bu.edu/psa/wd-example.htm - [Cached](#) - [Similar](#) - [Print](#) [X](#)

### The mutated **subsequence** problem and locating conserved genes ...

by Ht. Chan - 2005 - Cited by 6 - Related articles

(1999): A conserved gene rarely comprises the same entire **sequence** in the two .... hot spots where a number of mutations cluster together in the same **region**. .... **Backbone**. Find the MWCS of A and B. We call this **subsequence** as the ... We **repeat** such refinement until no more problematic pairs of intervals are left. ...

bioinformatics.oxfordjournals.org/cgi/content/full/21/10/2271 - [Similar](#) - [Print](#) [X](#)

### New Approximation Algorithms for Longest Common Subsequences ...

by L. Bergroth - 1996 - Cited by 14 - Related articles

identical **sequence** of (not necessarily contiguous) symbols found in both strings. .... There's no guarantee that points on the **backbone** belong ... which define the **region** to be studied, all symbols of  $\Sigma^*$  .... length string after which the source begins to **repeat** the generated **sequence** is short, the dependencies ...

ieeexplore.ieee.org/iel4/5792/15455/00712960.pdf?arnumber=712980 - [Print](#) [X](#)

### Key to acedb graphic **display**

Mar 14, 2001 ... Below is an image of a **sequence display** for the AGR database. ... of the page to see a zoomed in view of just that **subsequence**. You can then use the 'Zoom out.' button to see the **region** of **sequence** surrounding your gene of interest ... **Backbone** marker, Single position or Best of Multiple positions. ...

ukcrop.net/agr/sequence\_display\_key - [Cached](#) - [Similar](#) - [Print](#) [X](#)

### Structural stability of short **subsequences** of the tropomyosin chain

featuring a pseudo-repeating heptad of amino acids ... the near C-terminal253-280 **subsequence**); a sub- **sequence**-particularly rich in potential intrachain .... clude that the contribution to the **backbone region** ...

doi.wiley.com/10.1002/bip.360350113 - [Print](#) [X](#)

### Site-Specific Phosphorylation of Lys-Ser-Pro **Repeat** Peptides from ...

by P Sharma - 1998 - Cited by 20 - Related articles - All 5 versions




Peptides 1 and 2 were derived from the NF-H **sequence** of mouse and human species, respectively, while peptides 3 and 4 were from the KSP **repeat region** of mouse ... and the listed **sequence**; mult is the number of additional times a **subsequence** occurs .... Figure 5 (a, top) Overlay of the **backbone** atoms of the 24 best ...

pubs.acs.org/cgi-bin/jtext?bichaw/37/i14/abs/bi972746o - [Print](#) [X](#)

### Mauve: Multiple Alignment of Conserved Genomic **Sequence** With ...




by ACE Darling - 2004 - Cited by 219 - Related articles - All 21 versions




... been developed to **display** the rearrangement structure of several genome sequences. .... Anchoring its alignment in unique **subsequences** provides Mauve with ... When scored

only on **regions** considered **backbone sequence**, the accuracy is .... **Repeat analysis** using tools such as **Repeat Masker**, **REPuter** (Kurtz et al. ...  
[genome.cshlp.org/content/14/7/11394.full](http://genome.cshlp.org/content/14/7/11394.full) -   

### Conservation of genetic information




by L.F. Harris - Cited by 10 - Related articles - All 11 versions

The **regions** of nucleotide **subsequence** similarity to the GRE in the GR DBD occur .... Therefore, we proposed that the zinc finger may serve as a structural **backbone** for ... critical regulatory element within the MMTV 5' long terminal **repeat**, ... **regions** which showed genetic **sequence** similarity to the GR DBD exons 3, ...  
[www.mst.umn.edu/Projects/mg90601/pnas1.html](http://www.mst.umn.edu/Projects/mg90601/pnas1.html) - [Cached](#) - [Similar](#) -   

**Comprehensive sequence analysis of the 182 predicted open reading ...**  
 [PubMed]; Holm L, Sander C. Database algorithm for generating protein **backbone** and **side-chain** ... Intragenic **repeat** structure and identification of a family of ... Identification of common molecular **subsequences**. J Mol Biol. ... [PubMed]; Sor F, Chéret G, Fabre F, Faye G, Fukuhara H. **Sequence of the HMR region on** ...  
[www.ncbi.nlm.nih.gov/pmc/articles/PMC2142145/](http://www.ncbi.nlm.nih.gov/pmc/articles/PMC2142145/) -   

### Cell - The SV40 72 bp **repeat** preferentially potentiates ...

by C. Kanduri - 2000 - Cited by 85 - Related articles

... in Simian Virus 40 (SV40) and various **sequence** mutants and insertion derivatives. ... Hence, stable curvature in this **region** of the VP1 gene is due to the five ... within a **region** of the helix **backbone** that is intrinsically curved, ... VRE(IFN $\alpha$ ), tetrameric repA, and tetrameric GAAAGT (a **subsequence** of repB) or ...  
[www.cell.com/abstract/S0092-8674\(83\)90470-1](http://www.cell.com/abstract/S0092-8674(83)90470-1) - [Cached](#) -   

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